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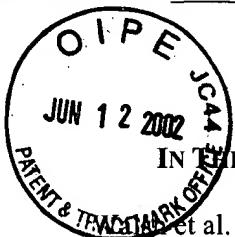
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Attorney's Docket No. 35052/204373(5052-53)

JUN 24 2002

PATENT

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IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

In re:
Appl No.: 09/689,430
Filed: October 12, 2000
For: ADENO-ASSOCIATED VIRUS VECTORS ENCODING
FACTOR VIII AND METHODS OF USING THE SAME

Confirmation No.: 7095
Group Art Unit: 1632
Examiner: Q. Li

June 12, 2002

Commissioner for Patents
Washington, DC 20231

SUBMITTAL OF FORMAL DRAWINGS
37 CFR § 1.85(c)

Sir:

In response to the requirement for new drawings as set forth in Paper No. 8 in the above application, there is enclosed herewith one set (25 sheets) of new formal drawings. It is requested that these new drawings be substituted for the originally filed drawings.

Respectfully submitted,

A handwritten signature in cursive ink that appears to read "Kathryn L. Coulter".

Kathryn L. Coulter
Registration No. 45,889

<p>Customer No. 00826 ALSTON & BIRD LLP Bank of America Plaza 101 South Tryon Street, Suite 4000 Charlotte, NC 28280-4000 Tel Raleigh Office (919) 862-2200 Fax Raleigh Office (919) 862-2260</p>	<p>CERTIFICATE OF EXPRESS MAILING "Express Mail" Mailing Label Number EL868643359US Date of Deposit: June 12, 2002 I hereby certify that this paper or fee is being deposited with the United States Postal Service "Express Mail Post Office to Addressee" service under 37 CFR 1.10 on the date indicated above and is addressed to: Commissioner for Patents, Washington, DC 20231.</p>
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<p>Nora C. Martinez</p>

FIG. 1A.

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10	20	30	40	50	
1234567890	1234567890	1234567890	1234567890	1234567890	
TGGCCACTOC	CCTCTCTGCGC	GCTCGCTCGC	TCACTGAGGC	CGGGCGACCA	50
AAGGTGCGCC	GAOGCGCGGG	CTTTGCGCGG	CGGGCGTCAG	TGAGCGAGCG	100
AGCGCGCAGA	GAGGGAGTGG	CCAACCTCCAT	CACTAGGGGT	TOCTCAGATC	150
TCTTCTAAAG	TAAACAGTAC	ATGAACTTT	ACCCCGITGC	TOGGCAAOGG	200
CCIGGCTCTG	GOCAAGTGTG	TGCTGACGCA	ACCCCGACTG	GCTGGGGCTT	250
GGCCATAGGC	CATCAGCGCA	TGCGGATCTC	AGTGTGGTT	TGCAAGAGGA	300
AGCAAAAAGC	CTCTCCACCC	AGGCGTGGAA	TGTTTCCACC	CAATGCGAG	350
CAGTGTTGTT	TTGCAAGAGG	AAGCAAAAAG	CCCTCTCCACC	CAGGCGTGG	400
CTCGAGAGCT	TOGACCACCA	TGCAAATAGA	GCTCTCCACC	TGCTTCCTTC	450
M etGlnIleG1	uLeuSerThr	CysPhePheL			
TGIGCCCTTT	GCGATTCTGC	TTTAGTGCCTA	CCAGAAAGATA	CTACCTGGGT	500
euCysLeuLe	uArgPheCys	PheSerAlaT	hrArgArgTy	rTyrLeuGly	
GCAGTGGAAC	TGTCATGGGA	GTATATGCAA	AGTGTCTCG	GTGAGCTGCC	550
AlaValGluL	euSerTrpAs	pTyrMetGln	SerAspLeuG	lyGluLeuPr	
TGIGGAAOGCA	AGATTTCCTC	CTAGAGTGCCT	AAAATCTTT	CCATTCAACA	600
oValAspAla	ArgPheProP	roArgValPr	oLysSerPhe	ProPheAsnT	
CCCTCAGTCGT	GTACAAAAAG	ACTCTGTTTG	TAGAATTCCAC	GGITCACCTT	650
hrSerValVa	1TyrLysLys	ThrLeuPheV	alGluPheTh	rValHisLeu	
TTCAACATCG	CTAACGCAAG	GCACACCTGG	ATGGGCTCTG	TAGGTCTTAC	700
PheAsnIleA	laLysProAr	gProProTrp	MetGlyLeuL	euGlyProTh	
CATCCAGGCT	GAGGTTTAIG	ATACAGTGGT	CATTACACTT	AAGAACATGG	750
rIleGlnAla	GluValTyrA	spThrValVa	lIleThrLeu	LysAsnMetA	
CTTCCCCATCC	TGTCAGCTTT	CATGCTGTTG	GTGTATCTA	CTGGAAAGCT	800
laSerHisPr	oValSerLeu	HisAlaValG	lyValSerTy	rTrpLysAla	
TCTGAGGGAG	CTGAATATGA	TGATCAGACCC	AGTCAAAGGG	AGAAAGAAGA	850
SerGluGlyA	laGluTyrAs	pAspGlnThr	SerGlnArgG	luLysGluAs	
TGATAAAAGTC	TTCCTGGTG	CAAGCCATAC	ATATGCTGG	CAGGTCTGAA	900
pAspLysVal	PheProGlyG	lySerHisTh	rTyrValTrp	GlnValLeuL	
AAGAGAAATGG	TOCAAATGCC	TCTGACCCAC	TGTGCTTAC	CTACTCATAT	950
ysGluAsnGl	yProMetAla	SerAspProL	euCysLeuTh	rTyrSerTyr	

02/24/2013

FIG. 1B.

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10	20	30	40	50	
1234567890	1234567890	1234567890	1234567890	1234567890	
CTTTCTCATG	TGGACCTGGT	AAAAGACTTG	AATTCAAGGCC	TCATGGAGC	1000
IeuSerHisV	aLAspLeuVa	LLysAspLeu	AsnSerGlyL	eUileGlyAl	
CCTACTAGTA	TGTAGAGAAAG	GGAGCTCTGGC	CAAGGAAAAG	ACACAGACCT	1050
aleuLeuVal	CysArgGluG	LySerLeuAl	aLysGluLys	ThrGlnThrL	
TGCACAAATT	TATACTACTT	TTTGCTGTAT	TTGATGAAGG	GAAAAGTGG	1100
euHisLysPh	eIleLeuLeu	PheAlaValP	heAspGluGl	yLysSerTrp	
CACTCAGAAA	CAAAGAACTC	CTTGATGCCAG	GATAGGGATG	CTGCATCTGC	1150
HisSerGluT	hrLysAsnSe	rLeuMetGln	AspArgAspA	laAlaSerAl	
TCGGGCTTGG	CCCTAAAATGC	ACACAGICAA	TGGTTATGTA	AAACAGGTCIC	1200
aArgAlaTrp	ProLysMetH	isThrValAs	nGlyTyrVal	AsnArgSerL	
TGOCAGGTCT	GATTGGATGC	CACAGGAAAT	CAGCTATTG	GCATGIGATT	1250
euProGlyLe	uIleGlyCys	HisArgLySS	erValTyrTr	pHisValIle	
GGAATGGGCA	CCACTCTGTA	AGTGCACITCA	ATATTCTCG	AAGGTACAC	1300
GlyMetGlyT	hrThrProGl	uValHisSer	IlePheLeuG	luGlyHisTh	
ATTCTCTGIG	AGGAAOCATC	GOCAAGGOGTC	CTTGGAAATC	TOGCCAATAA	1350
rPheLeuVal	ArgAsnHisA	rgGlnAlaSe	rLeuGluIle	SerProIleT	
CTTTCCTTAC	TGCTCAAACA	CTCTTGATGG	ACCTTGGACA	GTTTCTACTG	1400
hrPheLeuTh	rAlaGlnThr	LeuLeuMetA	spLeuGlyGl	nPheLeuLeu	
TTTTGTCATA	TCTCTTCCOA	CCAACATGAT	GGCATGGAAG	CTTATGICAA	1450
PheCysHisI	leSerSerHi	sGlnHisAsp	GlyMetGluA	laTyrValLy	
AGTAGACAGC	TGTCCAGAGG	AAOCCCCACT	AOGAATGAAA	AATAATGAAG	1500
sValAspSer	CysProGluG	luProGlnLe	uArgMetLys	AsnAsnGluG	
AAGCGGAAGA	CTATGATGAT	GATCTTACTG	ATTCTGAAAT	GGATGTGGTC	1550
luAlaGluAs	pTyrAspAsp	AspLeuThrA	spSerGluMe	tAspValVal	
AGGTTTGAATG	ATGACAACTC	TCCTTCTTT	ATCCAAATTC	GCTCAGTTC	1600
ArgPheAspA	spAspAsnSe	rProSerPhe	IleGlnIleA	rgSerValAl	
CAAGAACCAT	CCTAAAACCTT	GGGTACATTA	CATTGCTGCT	GAAGAGGAGG	1650
aLysLysHis	ProLysThrT	rpValHisTy	rIleAlaAla	GluGluGluA	
ACTGGGACTA	TGCTCCCTTA	GTCCTCGGCC	COGATGACAG	AAGTTATAAA	1700
spTrpAspTy	rAlaProLeu	ValLeuAlaP	roAspAspAr	gSerTyrLys	
AGTCAATATT	TGAACAATGG	CCCTCAGOGG	ATTGGTAGGA	AGTACAAAAAA	1750
SerGlnTyrL	euAsnAsnGl	yProGlnArg	IleGlyArgL	ysTyrLysLy	
AGTCGGATT	ATGGCATAACA	CAGATGAAAC	CTTAAAGACT	CGTGAAGCTA	1800
sValArgPhe	MetAlaTyrT	hrAspGluTh	rPheLysThr	ArgGluAlaI	
TTCAGCATGA	ATCAGGAATC	TTGGGACCTT	TACTTTATGG	GGAAGTGG	1850
leGlnHisGl	uSerGlyIle	LeuGlyProL	euLeuTyrGl	yGluValGly	
GACACACTGT	TGATTATATT	TAAGAAATCAA	GCAAGCAGAC	CATATAACAT	1900
AspThrLeuL	euIleIlePh	eLysAsnGln	AlaSerArgP	roTyrAsnIl	

FIG. 1C.

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10	20	30	40	50	
1234567890	1234567890	1234567890	1234567890	1234567890	
CTACCCCTCAC	GGAATCACTG	ATGTCGGTCC	TTTGTATTC	AGGAGATTAC	1950
eTyrProHis	GlyIleThrA	spValArgPr	oleuTyrSer	ArgArgLeuP	
CAAAAGGTGT	AAAACATTTC	AAGGATTTTC	CAATTCTGCC	AGGAGAAATA	2000
roLysGlyVa	lLysHisLeu	LysAspPheP	roIleLeuPr	oGlyGluIle	
TTCAAATATA	AATGGACAGT	GACTGTAGAA	GATGGGCCAA	CTAAATCAGA	2050
PheLysTyrL	ysTrpIhrVa	lThrValGlu	AspGlyProT	hrLysSerAs	
TOCTCGGTGC	CTGACCOOGCT	ATTACTCTAG	TTTGTAAAT	ATGGAGAGAG	2100
pProArgCys	LeuThrArgT	yrTyrSerSe	rPheValAsn	MetGluArgA	
ATCTAGCTTC	AGGACTCATT	GGCCCTCTCC	TCATCTGCTA	CAAAGAACCT	2150
spLeuAlaSe	rGlyLeuIle	GlyProLeuL	euIleCysTy	rLysGluSer	
GTAGATCAAA	GAGGAAACCA	GATAATGTCA	GACAAGAGGA	ATGTCATCCT	2200
ValAspGlnA	rgGlyAsnGl	nIleMetSer	AspLysArgA	srValIleLe	
GTTTCTGTIA	TTTGATGAGA	ACCGAAGCTG	GTACCTCAC	GAGAATATAC	2250
uPheSerVal	PheAspGluA	snArgSerTr	pTyrLeuThr	GluAsnIleG	
AAAGCTTTCT	CCCCAATCCA	GCTGGAGTGC	AGCTTGAGGA	TCCAGAGITC	2300
InArgPheLe	uProAsnPro	AlaGlyWalG	InLeuGluAs	pProGluPhe	
CAAGCTCTCA	ACATCATGCA	CAGCATCAAT	GGCTATGTTT	TIGATAGTTT	2350
GlnAlaSerA	snIleMetHi	sSerIleAsn	GlyTyrValP	heAspSerLe	
CCAGTTGTICA	GTTGTTTGC	ATGAGGTGGC	ATACTGGTAC	ATTCTAACGA	2400
uGlnLeuSer	ValCysLeuH	isGluValAl	aTyrTrpTyr	IleLeuSerI	
TTGGAGCACA	GACTGACTTC	CTTTCIGCT	TCTTCTCTGG	ATATACCTTC	2450
leGlyAlaGl	nThrAspPhe	LeuSerValP	hePheSerGl	yTyrThrPhe	
AAACACAAAA	TGGTCTATGA	AGACACACTC	ACCCATTTOC	CATTCTCAGG	2500
LysHisLysM	etValTyrGl	uAspThrLeu	ThrLeuPheP	roPheSerGl	
AGAAACCTGTC	TTCATGTGCA	TGGAAAACCC	AGGICTATGG	ATTCTGGGGT	2550
yGluThrVal	PheMetSerM	etGluAsnPr	oGlyLeuTrp	IleLeuGlyC	
GOCACAACTC	AGACTTTGG	AACAGAGGCA	TGACCGOCTT	ACTGAAGGTT	2600
ysHisAsnSe	rAspPheArg	AsnArgGlyM	etThrAlaLe	uLeuLysVal	
TCTAGTGTG	ACAAGAACAC	TGGTGATTAT	TACCGAGGACA	GTATGAAGA	2650
SerSerCysA	spLysAsnTh	rGlyAspTyr	TyrGluAspS	erTyrGluAs	
TATTCAGCA	TACITGCTGA	GTAAAAACAA	TGCCATTGAA	CCAAGAACCT	2700
pIleSerAla	TyrLeuLeuS	erLysAsnAs	nAlaIleGlu	ProArgSerP	
TCTCCCAGAA	TTCAAGACAC	OCTAGCACTA	GGCAAAAGCA	ATTTAATGOC	2750
heSerGlnAs	nSerArgHis	ProSerThrA	rgGlnLysGl	nPheAsnAla	
ACCCCAOCAG	TCTTGAACAG	CCATCAACGG	GAAATAACTC	GTACTACTCT	2800
ThrProProV	alleuLysAr	gHisGlnArg	GluIleThrA	rgThrThrLe	
TCAGTCAGAT	CAAGAGGAA	TIGACTATGA	TGATACCATA	TCAGTTGAAA	2850
uGlnSerAsp	GlnGluGluI	leAspTyrAs	pAspThrIle	SerValGluM	

FIG. 1D.

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10	20	30	40	50	
1234567890	1234567890	1234567890	1234567890	1234567890	
TGAAGAAGGA	AGATTTGAC	ATTATGATG	AGGATGAAAA	TCAGAGOC	2900
etLysLysGl	uAspPheAsp	IleTyrAspG	luAspGluAs	nGlnSerPro	
GGCAGCTTTC	AAAAGAAAAC	AOGACACTAT	TTTATTGCTG	CAGIGGAGAG	2950
ArgSerPheG	lnLysLysTh	rArgHisTyr	PheIleAlaA	laValGluAr	
GCTCTGGGAT	TATGGGATGA	GTAGCTOCCC	ACATGTTCTA	AGAAACAGGG	3000
gLeuTrpAsp	TyrGlyMetS	erSerSerPr	oHisValLeu	ArgAsnArgA	
CTCAGAGTGG	CAGTGTGCGT	CAGTTCAAGA	AAGTTGTTT	OCAGGAATT	3050
laGlnSerGl	ySerValPro	GlnPheLysl	ysValValPh	eGlnGluPhe	
ACTGATGGCT	OCTTACICA	GCCTTATAC	CGIGGAGAAC	TAAATGAACA	3100
ThrAspGlyS	erPheThrGl	nProLeuTyr	ArgGlyGluL	euAsnGluHi	
TTTGGGACTC	CTGGGGCCAT	ATATAAGAGC	AGAAGTGAA	GATAATATCA	3150
sLeuGlyLeu	LeuGlyProT	yrIleArgAl	aGluValGlu	AspAsnIleM	
TGGTAACITT	CAGAAATCG	GCCTCTCGIC	OCTATTCCIT	CTATTCTAGC	3200
etValThrPh	eArgAsnGln	AlaSerArgP	roTyrSerPh	eTyrSerSer	
CTTATTTCCT	ATGAGGAAGA	TCAGAGGCAA	GGAGCAGAAC	CTAGAAAAAA	3250
LeuIleSerT	yrGluGluAs	pGlnArgGln	GlyAlaGluP	roArgLysAs	
CTTGTCAAG	OCTAATGAAA	CCAAAACITA	CITTTGGAAA	GTGCAACATC	3300
nPheValLys	ProAsnGluT	hrLysThrTy	rPheTrpLys	ValGlnHisH	
ATATGGCACC	CACTAAAGAT	GAGTTTGACT	GCAAAGCCTG	GGCTTATTTC	3350
isMetAlaPr	oThrLysAsp	GluPheAspC	ysLysAlaTr	pAlaTyrPhe	
TCTGATGTIG	ACCTGGAAAAA	AGATGTGCCAC	TCAGGOCIGA	TTGGACCOCT	3400
SerAspValA	spLeuGluLy	sAspValHis	SerGlyLeuI	leGlyProLe	
TCTGGCTCTG	CACACTAACCA	CACTGAACCC	TGCTCATGGG	AGACAAGTGA	3450
uLeuValCys	HisThrAsnT	hrLeuAsnPr	oAlaHisGly	ArgGlnValT	
CAGTACAGGA	ATTTGCTCTG	TTTTTCACCA	TCTTTGATGA	GACCAAAAGC	3500
hrValGlnGl	uPheAlaLeu	PhePheThrI	lePheAspGl	uThrLysSer	
TGGTACTTCA	CTGAAAATAT	GGAAAGAAC	TGCAGGGCTC	CCTGCAATAT	3550
TrpTyrPheT	hrGluAsnMe	tGluArgAsn	CysArgAlaP	roCysAsnIl	
CCAGATGGAA	GATCCCACIT	TTAAAGAGAA	TTATOGCTTC	CATGCAATCA	3600
eGlnMetGlu	AspProThrP	heLysGluAs	nTyrArgPhe	HisAlaIleA	
ATGGCTACAT	AATGGATACA	CTACCTGGCT	TAGTAATGGC	TCAGGATCAA	3650
snGlyTyrIL	eMetAspThr	LeuProGlyL	euValMetAl	aGlnAspGln	
AGGATTCGAT	GGTATCTGCT	CAGCATGGCC	AGCAATGAAA	ACATCCATT	3700
ArgIleArgT	rpTyrLeuLe	uSerMetGly	SerAsnGluA	snIleHisSe	
TATTCATTTC	AGTGGACATG	TGTTCACTGT	ACGAAAAAAA	GAGGAGTATA	3750
rIleHisPhe	SerGlyHisV	alPheThrVa	1ArgLysLys	GluGluTyrL	
AAATGGCACT	GTACAATCTC	TATCCAGGIG	TTTTTGAGAC	AGTGGAAATG	3800
ysMetAlaLe	uTyrAsnLeu	TyrProGlyV	alPheGluTh	rValGluMet	

FIG. 1E.

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10	20	30	40	50	
1234567890	1234567890	1234567890	1234567890	1234567890	
TTACCATOCA	AAGCTGGAAT	TTGGCGGGTG	GAATGCCTTA	TTGGCGAGCA	3850
LeuProSerL	ysAlaGlyI1	eTrpArgVal	GluCysLeuI	IeGlyGluHi	
TCTACATGCT	GGGATGAGCA	CACTTTTCT	GGGTGACAGC	AATAAGTGTG	3900
sLeuHisAla	GlyMetSerT	hrLeuPheLe	uValTyrSer	AsnLysCysG	
AGACTCCCT	GGGAATGGCT	TCTGGACACA	TTAGAGATT	TCAGATTACA	3950
InThrProLe	uGlyMetAla	SerGlyHisI	IeArgAspPh	eGlnIleThr	
GCTTCAGGAC	AATATGGACA	GIGGGOOCCA	AACCTGGOCA	GACITCATT	4000
AlaSerGlyG	InTyrGlyGl	nTrpAlaPro	LysLeuAlaA	rgLeuHisTy	
TTCGGGATCA	ATCAAATGCT	GGAGCACCAA	GGAGGCCCTT	TCTTGGATCA	4050
rSerGlySer	IleAsnAlaT	rpSerThrLy	sGluProPhe	SerTrpIleL	
AGGTGGATCT	GTTGGCAACCA	ATGATTATTC	ACGGCATCAA	GACCCAGGGT	4100
ysValAspLe	uLeuAlaPro	MetIleIleH	isGlyIleLy	sThrGlnGly	
GOOGTCAGA	AGTCTCTCCAG	CCCTCACATC	TCTCAGTTA	TCAICATGTA	4150
AlaArgGlnL	ysPheSerSe	rLeuTyrIle	SerGlnPheI	IeIleMetTy	
TAGCTTGAT	GGGAAGAAGT	GGCAGACTTA	TCGAGGAAAT	TOCACTGGAA	4200
rSerLeuAsp	GlyLysLysT	rpGlnThrTy	rArgGlyAsn	SerThrGlyT	
CCTTAATGGT	CTTCTTTGGC	AATGTGGATT	CATCTGGAT	AAAACACAAT	4250
hrLeuMetVa	lPhePheGly	AsnValAspS	erSerGlyI1	eLysHisAsn	
ATTTTAAC	CTCCAATTAT	TGCTCGATAC	ATCCGTTTGC	ACCCAACCTCA	4300
IlePheAsnP	roProIleI1	eAlaArgTyr	IleArgLeuH	isProThrHi	
TTATAGCATT	OGCAGCACTC	TTGGCATGGA	GITGATGGC	TGIGATTAA	4350
sTyrSerIle	ArgSerThrL	euArgMetGl	uLeuMetGly	CysAspLeuA	
ATAGITGCAG	CATGOCATTG	GGAATGGAGA	GTAAAGCAAT	ATCAGATGCA	4400
snSerCysSe	rMetProLeu	GlyMetGluS	erLysAlaI1	eSerAspAla	
CAGATTACTG	CTTCATCTA	CTTTACCAAT	ATGTTTGCAC	CTGGCTCTOC	4450
GlnIleThrA	laSerSerTy	rPheThrAsn	MetPheAlaT	hrTrpSerPr	
TTCAAAAGCT	CGACTTCACC	TCCAAGGGAG	GAGTAATGCC	TGGAGACCTC	4500
oSerLysAla	ArgLeuHisL	euGlnGlyAr	gSerAsnAla	TrpArgProG	
AGGTGAATAA	TCCAAAAGAG	TGGCTGCAAG	TGGACTTCCA	GAAGACAATG	4550
InValAsnAs	nProLysGlu	TrpLeuGlnV	alAspPheGl	nLysThrMet	
AAAGTCACAG	GAGTAACTAC	TCAGGGAGTA	AAATCTCTGC	TTACCAGCAT	4600
LysValThrG	IyValThrTh	rGlnGlyVal	LysSerLeuL	euThrSerMe	
GTATGTGAAG	GAGTTOCTCA	TCTCCAGCAG	TCAAGATGGC	CATCAGTGG	4650
tTyrValLys	GluPheLeuI	IeSerSerSe	rGlnAspGly	HisGlnTrpT	
CTCTCTTTT	TCAGAATGGC	AAAGTAAAGG	TTTTTCAGGG	AAATCAAGAC	4700
hrLeuPhePh	eGlnAsnGly	LysValLysV	alPheGlnGl	yAsnGlnAsp	
TOCTTCACAC	CTGIGGTGAA	CCTCTAGAC	OCACCGTTAC	TGACTCGCTA	4750
SerPheThrP	roValValAs	nSerLeuAsp	ProProLeuL	euThrArgTy	

FIG. 1F.

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10	20	30	40	50	
1234567890	1234567890	1234567890	1234567890	1234567890	
OCTTCGAATT	CACCCCCAGA	GTTGGGTGCA	CCAGATTGCC	CTGAGGGATGG	4800
rLeuArgIle	HisProGlnS	erTrpValHi	sGlnIleAla	LeuArgMetG	
AGGTTCTGGG	CTGCGAGGCCA	CAGGACCTCT	ACTGACTCGA	GOGAGTTCTT	4850
luValLeuGl	yCysGluAla	GlnAspLeuT	yr...		
CTGAGGGGAT	CGGCAATAAA	AAGACAGAAT	AAAACGCAOG	GGTGTGGGT	4900
CGTTTGTTCG	GATCCAGATC	TAGGAACCCC	TAGTGATGGA	GTTGGCCACT	4950
CCCTCTCTGC	GOGCTCGCTC	GCTCACTGAG	GGGGGGGGGG	CAAAGCCCGG	5000
GCGTGGGCG	ACCTTGGTC	GGCCGGCGTC	AGTGAGCGAG	GGAGGCGCGA	5050
GAGAGGGAGT	GGCCAACCCCC	CCCCCCCCCCC	CCCGTGCAGC	CCAGCTGCAT	5100
TAATGAATCG	GCACACGCGC	GGGGAGAGGC	GGTTTGGTAA	TTGGGCGCTC	5150
<u>TTCGGCTTCC</u>	<u>TOGCTCACTG</u>	<u>ACTCGCTGCG</u>	<u>CTGGTTCGTT</u>	<u>CGGCTGCGGC</u>	5200
<u>GAGGGTATC</u>	<u>AGCTCACTCA</u>	<u>AAGGGGTA</u>	<u>TACGGTTATC</u>	<u>CACAGAACATCA</u>	5250
<u>GGGGATAACG</u>	<u>CAGGAAAGAA</u>	<u>CATGTGAGCA</u>	<u>AAAGGCCAGC</u>	<u>AAAAGGCCAG</u>	5300
<u>GAACCGTAAA</u>	<u>AAGGCGCGT</u>	<u>TGCTGGGTT</u>	<u>TTTCCATAGG</u>	<u>CTCGCGCCOC</u>	5350
<u>CTGACGAGCA</u>	<u>TCACAAAAAT</u>	<u>CGACGCTCAA</u>	<u>GTCAGAGGTG</u>	<u>GCGAAACCG</u>	5400
<u>ACAGGACTAT</u>	<u>AAAGATAACCA</u>	<u>GGCGTTTCCC</u>	<u>CCTGGAAAGCT</u>	<u>CCCTCGTGC</u>	5450
<u>CTCTCCCTGT</u>	<u>CGGACCCCTGC</u>	<u>CGCTTACCGG</u>	<u>ATACCTGTCC</u>	<u>GCCTTTCTCC</u>	5500
<u>CTTCGGGAAG</u>	<u>CGTGGCGCTT</u>	<u>TCTCAATGCT</u>	<u>CACGCTGTAG</u>	<u>GTATCTCAGT</u>	5550
<u>TOGGTGTAGG</u>	<u>TCGTTCGCTC</u>	<u>CAAGCTGGC</u>	<u>TGTGTGGCACG</u>	<u>AACCCCGCGT</u>	5600
<u>TCAGCGCGAC</u>	<u>CGCTGCGCCT</u>	<u>TATCCGGTAA</u>	<u>CTATCGCTT</u>	<u>GAGTCGAACC</u>	5650
<u>CGGTAAAGACA</u>	<u>CGACTTATCG</u>	<u>CCACTGGCAG</u>	<u>CAGCCACTGG</u>	<u>TAACAGGATT</u>	5700

FIG. 1G.

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10	20	30	40	50	
1234567890	1234567890	1234567890	1234567890	1234567890	
<u>AGCAGAGOGA</u>	<u>GGTATGTAGG</u>	<u>CGGTGCTACA</u>	<u>GAGTTCTTGA</u>	<u>AGTGGTGGCC</u>	5750
<u>TAAC TAC CGGC</u>	<u>TAC ACT AGAA</u>	<u>GGAC AGT ATT</u>	<u>TGGT ATCT GC</u>	<u>GCT CTG CTGA</u>	5800
<u>AGCC AGTT AC</u>	<u>CIT CCG AAAA</u>	<u>AGAG TTGG TA</u>	<u>GCT CTG ATC</u>	<u>CGG CAA ACA A</u>	5850
<u>ACCA ACG CTG</u>	<u>GTAG CGG GTGG</u>	<u>TTTTTT GTT</u>	<u>TGCA ACC AGC</u>	<u>AGATTAC CG</u>	5900
<u>CAG AAAAAA AA</u>	<u>GGAT CTCA AG</u>	<u>AAG ATC TTT</u>	<u>GAT CTT TCT</u>	<u>ACGG GGT CTG</u>	5950
<u>ACG CT CAG TGT</u>	<u>GAAC GAAA AC</u>	<u>TCAC GTT AAG</u>	<u>GGAT TTT GGT</u>	<u>CAT GAG ATTA</u>	6000
<u>TCA AAA AGGA</u>	<u>TCT TAC CTA</u>	<u>GAT CCT TTT A</u>	<u>AAT TAAA AT</u>	<u>GAAG TTT TAA</u>	6050
<u>AT CAAT CTAA</u>	<u>AGT ATAT ATG</u>	<u>AGT AAAC ITG</u>	<u>GTCT GAC AGT</u>	<u>TAC CAAT GCT</u>	6100
ueL...siHo	rPlaV...gr	AueLreSgrA	psAelIulGn	sAteMprTue	y1GelIres
GTG GCT GAC	TOCC COG TOGT	GTAGATAACT	ACGATA CGGG	AGGG CTT ACC	6150
InlGgrAlaV	y1GgrAgrAr	hTreSueL..	.reSlavOrP	orPreSlaVt	6200
ATCTGGCCCC	AGTG CTGCAA	TGATA CGCG	AGACCCACCG	TCAC CGGC TC	6250
eMn1GylGpr	TsiHn1GueL	reSlavAlAu	eLy1GlaVre	SlaVorPulG	6300
CAG ATT TATC	AGCA ATA AAC	CAG CCAG CG	GAAG GCG CGA	GCG CAG AAGT	6350
ueLnsAelIu	eLueLueLyl	GalAueLgrA	ehPorPgrAa	1AsyCehPsi	6400
GGT CCT GCAA	CTTT ATCCGC	CTCC ATCCAG	TCT ATT AATT	GTG CGGG GA	6450
HpsAn1GueL	syLeIigrAg	rAprTy1Grh	T.....nsA	nsAylGorPu	6500
AGCT AGAG TA	AGTAGT TCGC	CAG TTA ATAG	TTT GCG CAAC	GTG GTG OCA	6550
eL...ueLue	LryTnsAalA	ueL...ryTn	sAalAsyCgr	An1Gn1GprT	6600
TTG CTAC AGG	CAT CGG GIG	TCA CGCT CGT	CGTT GGT AT	GGCT TCAT TC	6650
n1G...ueLs	yCgrAorPrh	TlaVreSrht	rhTh1GryTo	rPsyLteM..	.
AGCT CGG GTT	CCC AAC GATC	AAG GCG AGT	ACAT GAT CCC	CCAT GTG GIG	.
.reSgrAnsA	y1GlaVelIu	eLalAueL..	.teMeIi y1G	prTrhTrhTs	.
CAAAAAGCG	GTT AGCT CCT	TCGGT CCT CC	GAT CGT GTG C	AGA AGT AAGT	.
yCehPueLor	P...reSgrA	grApsAulGr	eSgrAn1G..	.ehPryTrhT	.
TGG CGC AGT	GTT ATCA CT	ATGG TTAT GG	CAG CACT GCA	TAAT TCT CTT	.
orPgrAueLr	hTe l1laV..	.orP...orP	ueLlaValAr	yTnsAulG..	.
ACT GT CAT GC	CAT CGT AAG	ATG CTT TCT	GTG ACT GGT G	AGT ACT CAAC	.
.n1G...a1A	teMgrAueLe	1IreSsylnl	GreSnlGsiH	rhTreSueLp	.

FIG. 1H.

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10	20	30	40	50	
<u>1234567890</u>	<u>1234567890</u>	<u>1234567890</u>	<u>1234567890</u>	<u>1234567890</u>	
CAAGTCATTG	TGAGAATAGT	GTATGOGGCG	ACCGAGTTGC	TCTTGCOCGG	6700
rTrhTteMgr	AueLeIrhT	ryTalAalAl	aVreSnsAre	SsyLylGorP	
OGTCAATACG	GGATAATACC	GOGCCACATA	GCAGAACITTT	AAAAGTGCTC	6750
rhTueLlaVo	rPryTryTgr	AalAlaVryT	syCehPsyLu	eLueLala..	
ATCATTGGAA	AACGTTCTTC	GGGGGGAAAAA	CTCTCAAGGA	TCTTACCGCT	6800
....n1GehP	1aVnsAsyLo	rPalAehPla	VgrAueLreS	grAlaValAr	
GTTGAGATCC	AGITCGATGT	AACCCACTCG	TGCACCCAAC	TGATCTTCAG	6850
hTreSelIpr	TnsAreSrht	1aVprTulGs	iH1aVprTre	SelIsyLueL	
CATCTTTAC	TTTCACCAGC	GTTTCTGGGT	GAGCAAAAAC	AGGAAGGCCAA	6900
teMsyL...s	yL...prTgr	AsyLn1GrhT	ueLueLehPu	eLehPalAeh	
AATGOOGCAA	AAAAGGGAAT	AAGGGCGACA	CGGAAATGTT	GAATACTCAT	6950
PsiHgrAueL	ehPorPehPu	eLorPreSla	VreSelInsA	ehPlaV...1	
ACTCTTCCTT	TTTCAATATT	ATTGAAGCAT	TTATCAGGGT	TATTGTCICA	7000
TGAGCGGATA	CATATTGAA	TGTATTAGA	AAAATAAACAA	AATAGGGGTT	7050
CCCGGCACAT	TTCCOCGAAA	AGTGCACCT	GACGTCTAAG	AAACCATTAT	7100
TATCATGACA	TIAACCTATA	AAAATAGGCG	TATCACGAGG	COCTTTCGTC	7150
TGGGGGTTT	CGGTGATGAC	GGTGAAAACC	TCTGACACAT	GCAGCTCOOG	7200
GAGACGGTCA	CAGCTTGCT	GTAAGCGGAT	GGCGGGAGCA	GACAAGCG	7250
TCAGGGGGCG	TCAGCGGGTG	TTGGCGGGTG	TCGGGGCTGG	CTTAACATATG	7300
CGGCATCAGA	GCAGATTGTA	CTGAGAGTGC	ACCATATGCG	GTGTGAAATA	7350
CGGCACAGAT	GOGTAAGGAG	AAAATACCGC	ATCAGGAAAT	<u>TGTAAACGTT</u>	7400
<u>AATATTTTGT</u>	<u>TAAAATTGCG</u>	<u>GTTAAATTT</u>	<u>TGTAAATCA</u>	<u>GCTCATTTT</u>	7450
<u>TAACCAATAG</u>	<u>GGCGAAATCG</u>	<u>GCAAAATCCC</u>	<u>TTATAAATCA</u>	<u>AAAGAATAGA</u>	7500
<u>CGGAGATAGG</u>	<u>GTTGAGTGT</u>	<u>GTTCAGTTT</u>	<u>GGAACAAGAG</u>	<u>TOCCACTATTA</u>	7550
<u>AAGAACGTGG</u>	<u>ACTOCAAQGT</u>	<u>CAAAGGGCGA</u>	<u>AAAACCGTCT</u>	<u>ATCAGGGCGA</u>	7600

FIG. 11.

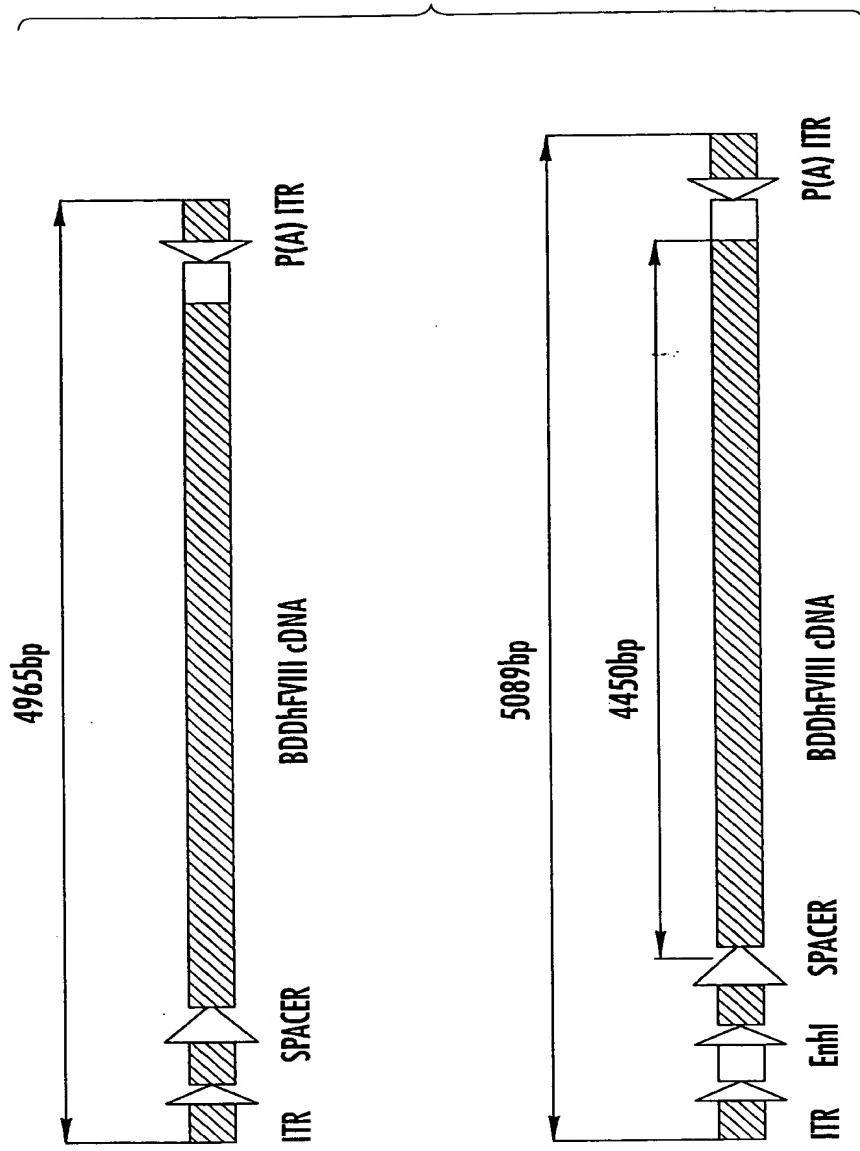
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10	20	30	40	50	
<u>1234567890</u>	<u>1234567890</u>	<u>1234567890</u>	<u>1234567890</u>	<u>1234567890</u>	
TGGCCCACTA	CGTGAACCAT	CACCTAAATC	AAGTTTTTG	GGGTCGAGGT	7650
<u>GGCGTAAAGC</u>	<u>ACTAAATCGG</u>	<u>AACCTAAAG</u>	<u>GGAGCCCCCG</u>	<u>ATTTAGAGCT</u>	7700
<u>TGAOGGGAA</u>	<u>AGCCCGCGAA</u>	<u>CGTGGCGAGA</u>	<u>AAGGAAGGGA</u>	<u>AGAAAGCGAA</u>	7750
<u>AGGAGCGGGC</u>	<u>GCTAGGGCGC</u>	<u>TGGCAAGTGT</u>	<u>AGGGTCAOG</u>	<u>CTGOGCGTAA</u>	7800
<u>CCAOCACACC</u>	<u>CGCCCGCGCTT</u>	<u>AATGCCCGGC</u>	<u>TACAGGGCGC</u>	<u>GTGCGGCCAT</u>	7850
<u>TCGOCATTCA</u>	<u>GGCTACGCAA</u>	<u>CTGTTGGAA</u>	<u>GGGCGATCGG</u>	<u>TGCGGGCGTC</u>	7900
<u>TTCGCTATTA</u>	<u>CGCCAGCTGG</u>	<u>CTGCAGGGGG</u>	<u>GGGGGGGGGG</u>	<u>GGGT</u>	7944

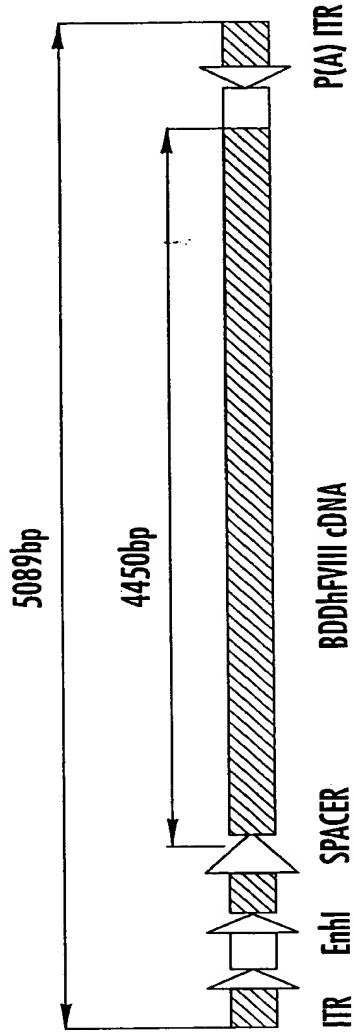
Title: Adeno-Associated Virus Vectors Encoding Factor VIII
and Methods of Using the Same
Inventor(s): Walsh et al.
Application No: 09/689,430
Atty Dkt No: 33052/204373(5052-53)

FIG. 2.

pDIZ2.



pDIZ6.



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Title: Adeno-Associated Virus Vectors Encoding Factor VIII
and Methods of Using the Same
Inventor(s): Walsh et al.
Application No: 09/689,430
Atty Dkt No: 35052/204373(5052-53)

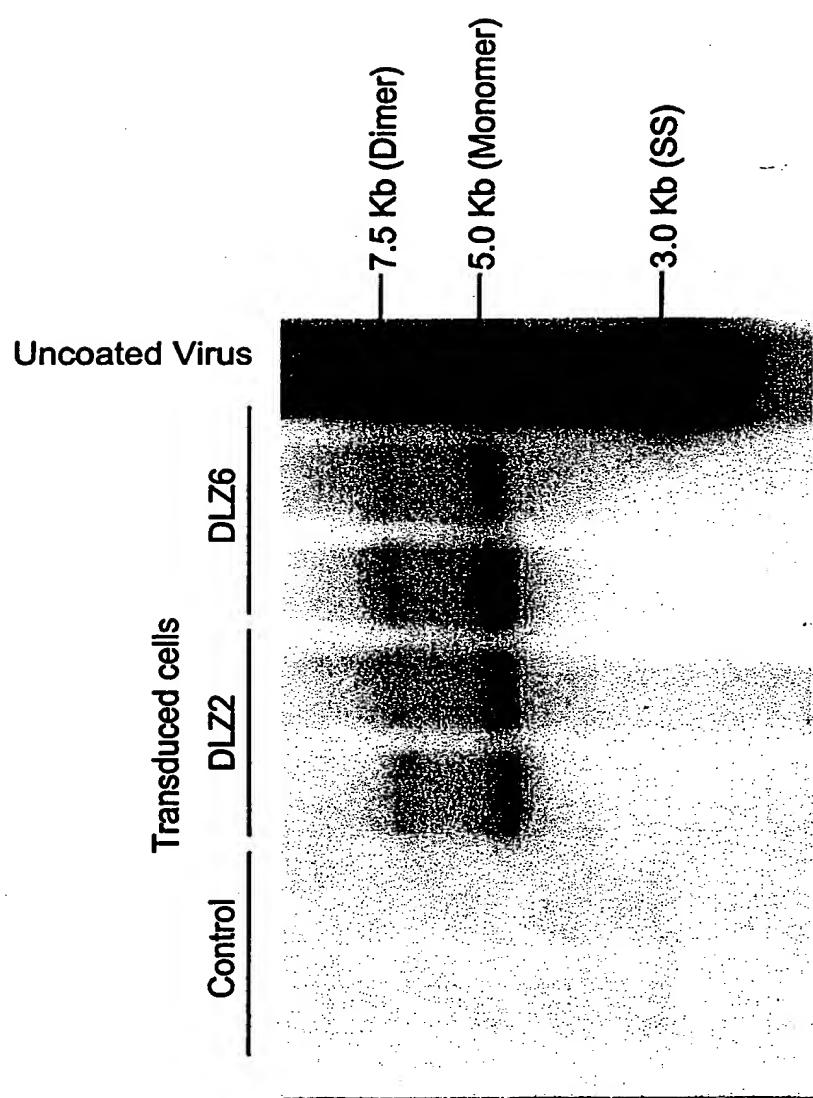
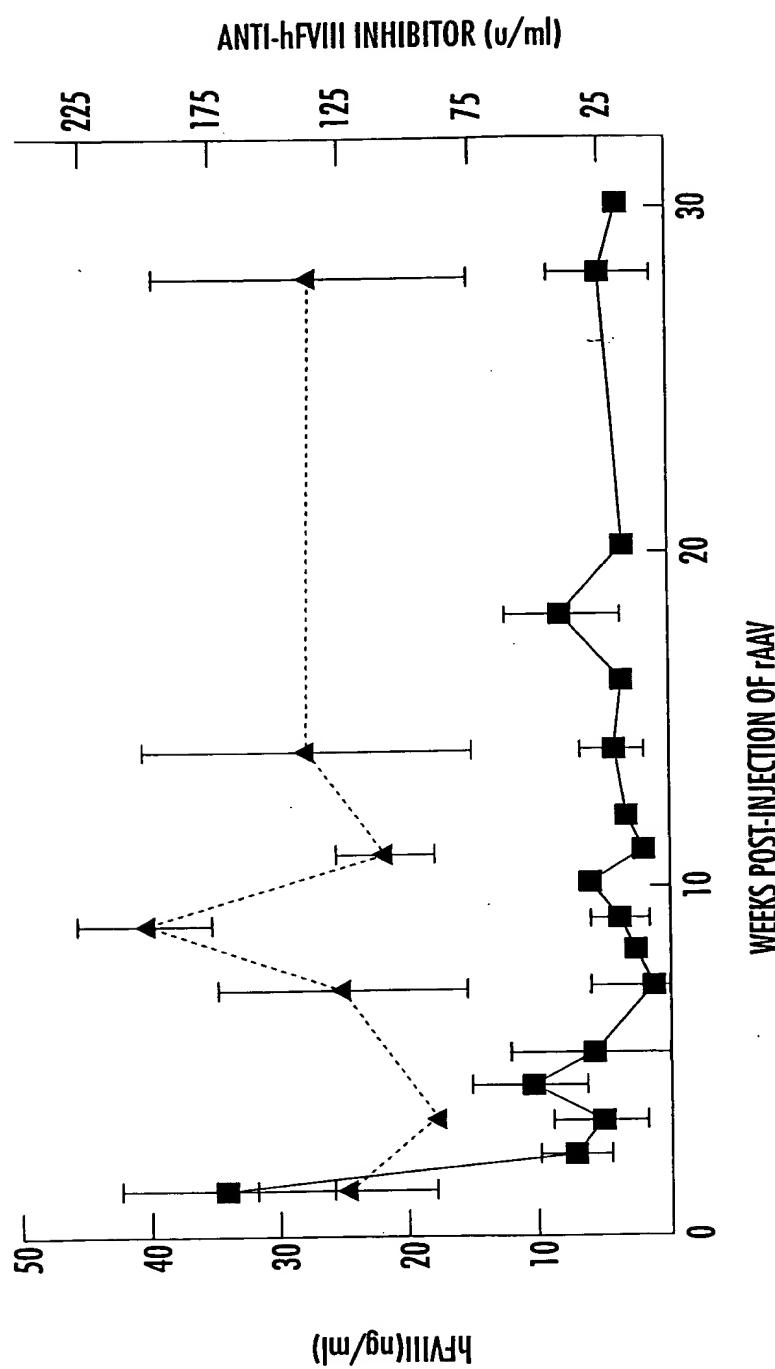


FIG. 3.

Title: Adeno-Associated Virus Vectors Encoding Factor VIII
and Methods of Using the Same
Inventor(s): Walsh et al.
Application No: 09/689,430
Atty Dkt No: 35052/204373(5052-53)

FIG. 4.A.



Title: Adeno-Associated Virus Vectors Encoding Factor VIII
and Methods of Using the Same
Inventor(s): Walsh et al.
Application No: 09/689,430
Atty Dkt No: 35052/204373(5052-53)

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FIG. 4.B.

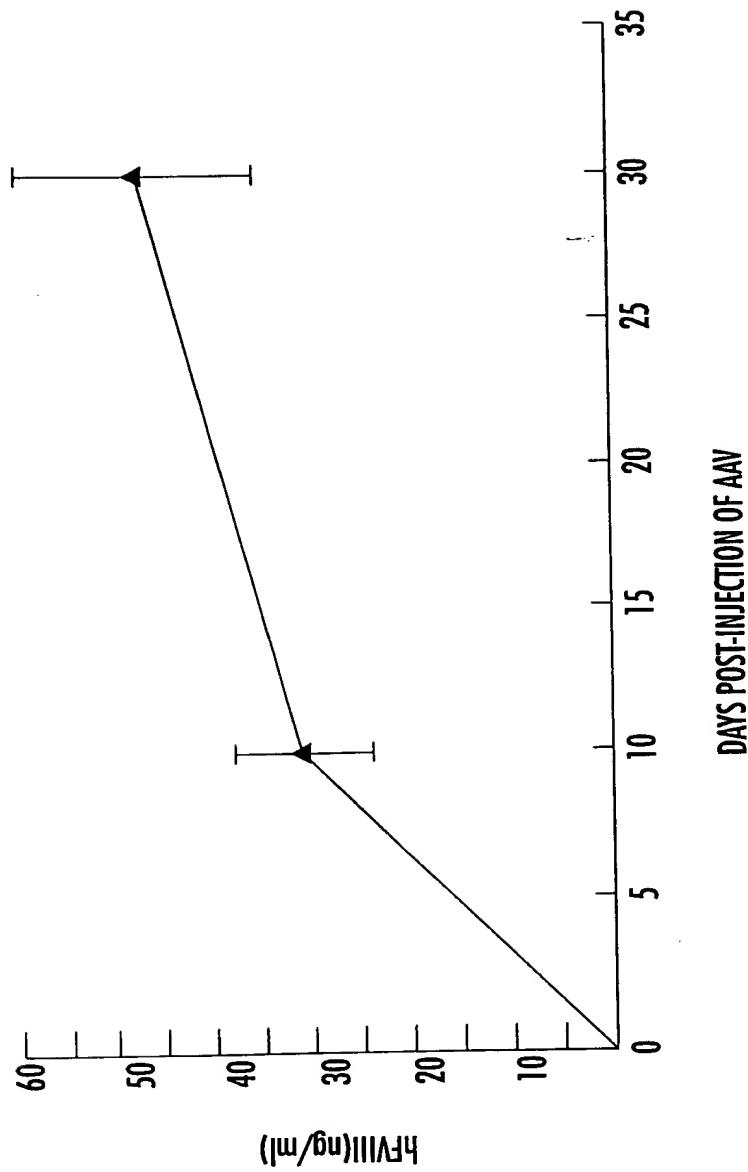


FIG. 5.A.

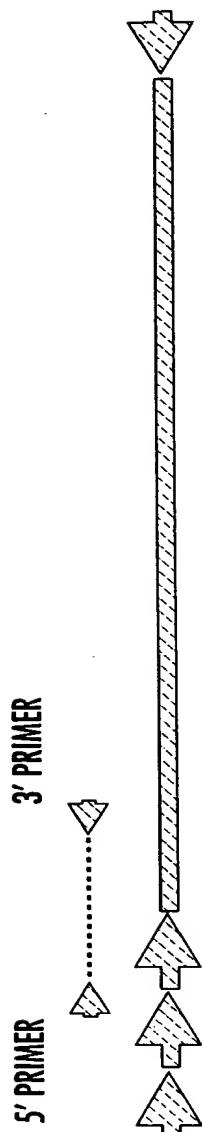
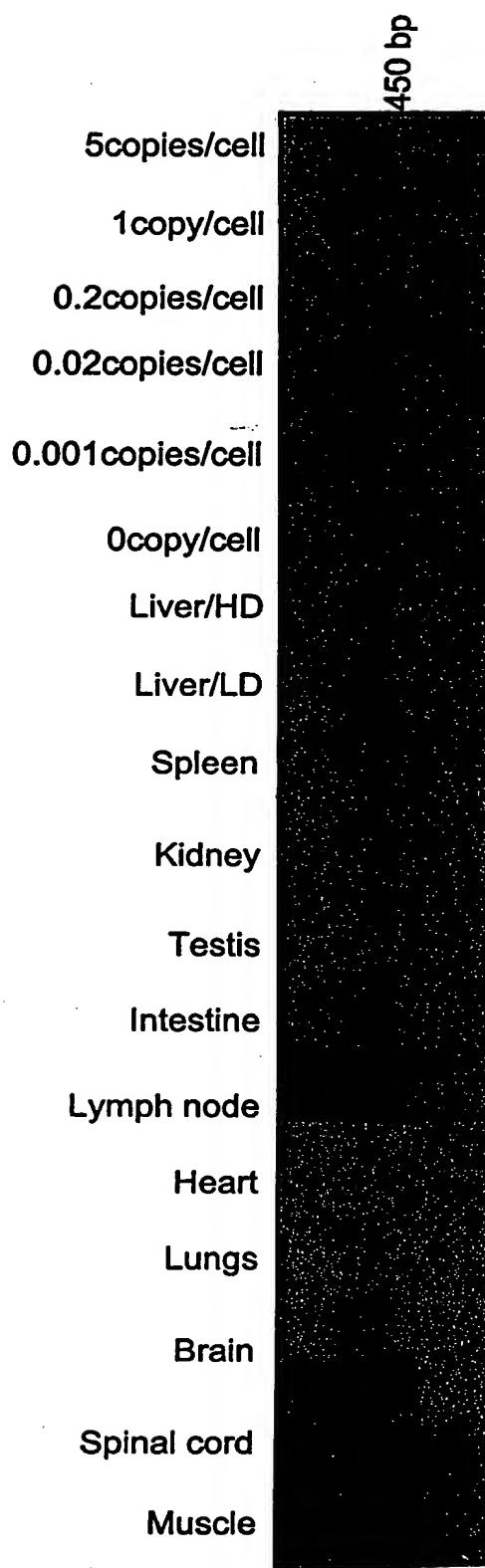


FIG. 5.B.

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Title: Adeno-Associated Virus Vectors Encoding Factor VIII
and Methods of Using the Same
Inventor(s): Walsh et al.
Application No: 09/689,430
Atty Dkt No: 35052/204373(5052-53)

FIG. 5.C.

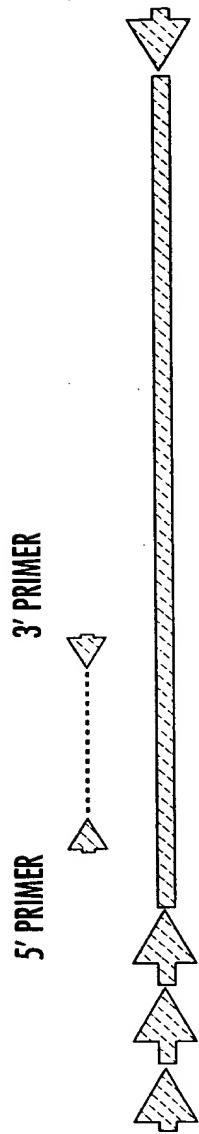


FIG. 5.D.

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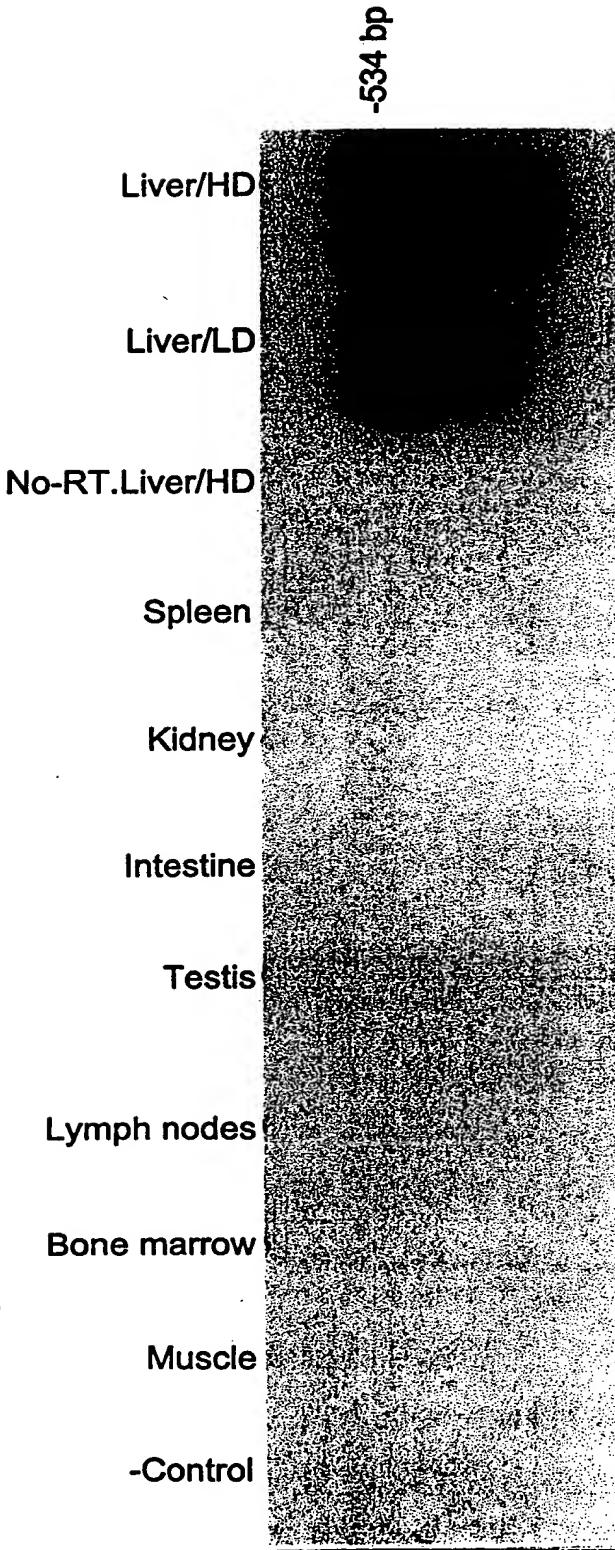


FIG. 5.E.

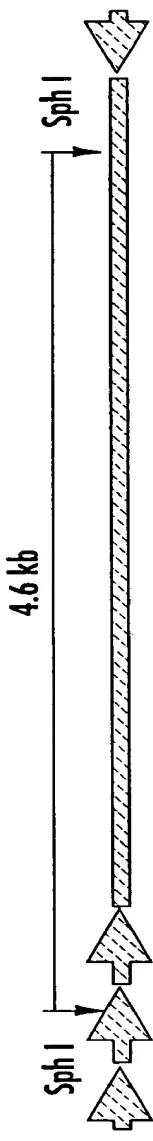


FIG. 5.F.

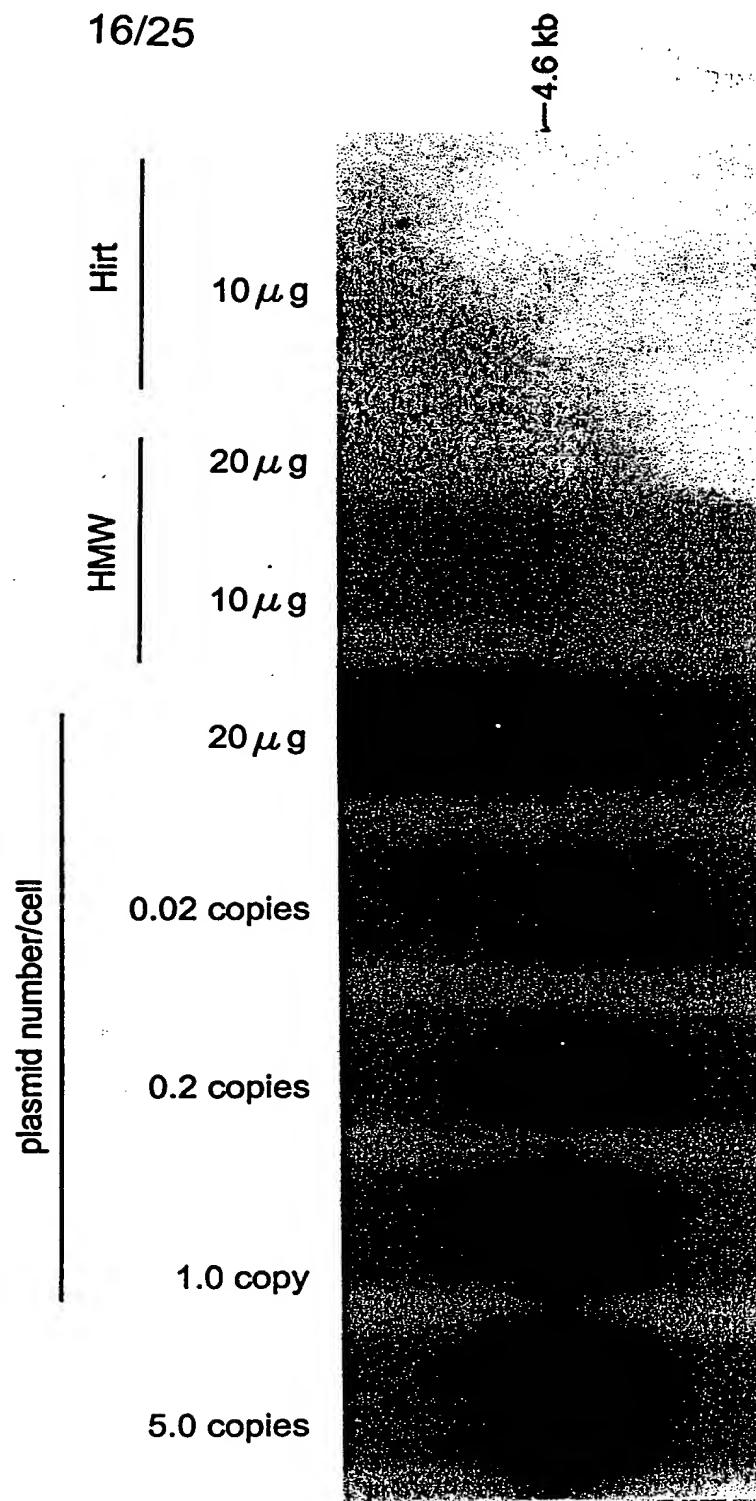


FIG. 6.A.

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10	20	30	40	50	
1234567890	1234567890	1234567890	1234567890	1234567890	
TGGCCACTGCC	CTCTCTGCGC	GCTGCTGCGC	TCACTGAGGC	GGGGCGACCA	50
AAGGTGCGCC	GAOGCCCCGGG	CITTGCGCGG	GGGGCGTCAG	TGAGCGAGCG	100
AGOGOGCAGA	GAGGGAGTGG	CCAACCTCCAT	CACTAGGGGT	TCCTCAGATC	150
TCTTCTAAG	TAAACAGTAC	ATGAACCTTT	ACCCCGTTGC	TGGGCAAOGG	200
CCTGGTCGT	GCCAAGTGT	TGCTGACGCA	ACCCOCACTG	GCTGGGGCTT	250
GGOCATAGGC	CATCAGCGCA	TGCGGATCTC	AGTGTGGTTT	TGCAAGAGGA	300
AGCAAAAAGC	CTCTCCACCC	AGGOCTGGAA	TGTTTCCACC	CAATGTGAG	350
CAGTGTGGTT	TTGCAAGAGG	AAGCAAAAAG	CCTCTCCACC	CAGGCGCTGGA	400
CTCGACCTCG	AGAGTACTTC	TAGAAATAAG	AGCCATGCAA	GTAGAGCTCT	450
				MetGln ValGluLeuT	
ACACCTGCTG	CTTTCTGIGC	CTTTGCGCT	TCAGCCCTAG	TGCCACCGAGA	500
yrThrCysCy	sPheLeuCys	LeuLeuProP	heSerLeuSe	rAlaThrArg	
AAATACTACC	TOGGTGCAGT	GGAACTGTOC	TGGGACTATA	TGCAAAGTIGA	550
LysTyrTyrL	euGlyAlaVa	1GluLeuSer	TrpAspTyrM	etGlnSerAs	
CCTGCTCAGT	GCGCTGCAOG	CGGATACAAG	CTTTCTTCC	AGGGTGCCAG	600
pLeuLeuSer	AlaLeuHisA	1aAspThrSe	rPheSerSer	ArgValProG	
GATCTTGCC	ACTCACCAOG	TCAGTCACGT	ACAGAAAGAC	TGTGTTTGT	650
lySerLeuPr	oLeuThrThr	SerValThrT	yrArgLysTh	rValPheVal	
GAGTTTACAG	ATGACCTTTT	CAACATTGCC	AAGGCCAGGC	CACCGTGGAT	700
GluPheThrA	spAspLeuPh	eAsnIleAla	LysProArgP	roProTrpMe	
GGGCGCTGCTG	GGTCCTACCA	TOCAGGCTGA	GGTTTATGAC	ACAGTGGTCA	750
tGlyLeuLeu	GlyProThrI	1eGlnAlaGl	uValTyrAsp	ThrValValI	
TTGTCCTTAA	GAACATGGCT	TCTCATCCTG	TCAGCCTTCA	CGCTGTTGGT	800
leValLeuLy	sAsnMetAla	SerHisProV	alSerLeuHi	sAlaValGly	
GTATCCTATT	GGAAAGCTTC	TGAAGGTGCT	GAGTATGAGG	ATCAGACCAAG	850
ValSerTyrT	rpLysAlaSe	rGluGlyAla	GluTyrGluA	spGlnThrSe	
CCAAAAGGAG	AAGGAAGATG	ATAATGTCAT	TOCTGGTGAAG	AGCCATAACCT	900
rGlnLysGlu	LysGluAspA	spAsnValII	eProGlyGlu	SerHisThrT	
ATGCTGGCA	GGTCCTGAAA	GAGAATGGCC	CAATGGCTC	TGATCCACCA	950
yrValTrpGl	nValLeuLys	GluAsnGlyP	roMetAlaSe	rAspProPro	

FIG. 6.B.

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10	20	30	40	50	
1234567890	1234567890	1234567890	1234567890	1234567890	
TGTCTCACCT	ACTCATATT	TTCACACGTG	GACCTGGIGA	AAGACCTGAA	1000
CysLeuThrT	yrSerTyrPh	eSerHisVal	AspLeuValL	ysAspLeuAs	
TTCAGGOCCTC	ATTGGAGGCC	TGCTGGTTTG	CAAAGAACGGG	AGCTCTGGCCA	1050
nSerGlyLeu	IleGlyAlaL	euLeuValCy	sLysGluGly	SerLeuAlaL	
AAGAAAGGAC	ACAGACCTTG	CAGGAATTG	TCCTACTTTT	TGCTGTATT	1100
ysGluArgTh	rGlnThrLeu	GlnGluPheV	alLeuLeuPh	eAlaValPhe	
GATGAAGGGA	AAAGTTGGCA	CTCAGAAACA	AATGOGICCTT	TGACACAGGC	1150
AspGluGlyL	ysSerTrpHi	sSerGluThr	AsnAlaSerL	euThrGlnAl	
TGAGGOCAG	CATGAGCTGC	ACAOCATCAA	TGGCTATGTA	AACAGGTCTC	1200
aGluAlaGln	HisGluLeuH	isThrIleAs	nGlyTyrVal	AsnArgSerL	
TGCGAGGCT	TACTGIGTGT	CACAAGAGAT	CAGCTTATTG	GCATGTGATT	1250
euProGlyLe	uThrValCys	HisLysArgS	erValTyrTr	pHisValIle	
GGAAATGGCCA	CCACCCCOGA	AGTGCCTCA	ATTTCCTCTCG	AAGGTACAC	1300
GlyMetGlyT	hrThrProGl	uValHisSer	IlePheLeuG	luGlyHisTh	
ATTCTTGIG	AGGAACCACC	GOCAAGCCTC	CTTGGAGATC	TCACCAATT	1350
rPheLeuVal	ArgAsnHisA	rgGlnAlaSe	rIeuGluIle	SerProIleT	
CTTCCTTAC	TGCTCAGACA	TTCTGATGG	ACCTTGGCCA	GTTTCTACTG	1400
hrPheLeuTh	rAlaGlnThr	PheLeuMetA	spLeuGlyG1	nPheLeuLeu	
TTTGTCTATA	TCCCTTOCCA	TCAACATGAT	GGTATGGAAG	CITATGTCAA	1450
PheCysHisI	leProSerHi	sGlnHisAsp	GlyMetGluA	laTyrValLy	
AGTAGATAGC	TGCCCCAGAGG	AAACCCCAGCT	GCGCATGAAA	AATAATGAAG	1500
sValAspSer	CysProGluG	luProGlnLe	uArgMetLys	AsnAsnGluA	
ATAAAAGATTA	TGATGATGGT	CITTTATGATT	CTGACATGGA	CGTAGITAGC	1550
spLysAspTy	rAspAspGly	LeuTyrAspS	erAspMetAs	pValValSer	
TTTGATGACG	ACAGCTCTC	TCCTTATAC	CAAATCOGCT	CAGTTGCCAA	1600
PheAspAspA	spSerSerSe	rProPheIle	GlnIleArgS	erValAlaLy	
GAAGCATCT	AAAACCTGGG	TCCACTATAT	TGCTGCTGAG	GAGGAGGACT	1650
sLysHisPro	LysThrTrpV	alHisTyrIl	eAlaAlaGlu	GluGluAspT	
GGGACTATGC	TOCCCTCAGGC	CCACCCCOCA	ATGATAGAAG	TCATAAAAAT	1700
rpAspTyrAl	aProSerGly	ProThrProA	snAspArgSe	rHisLysAsn	
CTGTATTGTA	ACAATGGTCC	TCAGCGGATT	GGTAAGAAGT	ACAAAAAAAGT	1750
LeuTyrLeuA	snAsnGlyPr	oGlnArgIle	GlyLysLysT	yrLysLysVa	
COGATTGIG	GCATACACAG	ATGAGACATT	TAAGACTCGT	GAAGCTATT	1800
lArgPheVal	AlaTyrThrA	spGluThrPh	eLysThrArg	GluAlaIleG	
AGTATGAATC	AGGAATCCIG	GGACCTTAC	TTTATGGAGA	AGTTGGAGAC	1850
InTyrGluSe	rGlyIleLeu	GlyProLeuL	euTyrGlyG1	uValGlyAsp	
ACACTGCTGA	TTATATTAA	GAATCAAGCC	AGCGGGCCAT	ATAACATCTA	1900
ThrLeuLeuI	leIlePheLy	sAsnGlnAla	SerArgProT	yrAsnIleTy	

FIG. 6.C.

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10	20	30	40	50	
1234567890	1234567890	1234567890	1234567890	1234567890	
CCCTCATGGG	ATCAATTATG	TCACTCCTCT	GCACACAGGG	AGATTGCCAA	1950
rProHisGly	IleAsnTyrV	alThrProLe	uHisThrGly	ArgLeuProL	
AAGGTGTGAA	ACATTGAAA	GATATGCCAA	TCTCTGOOGGG	AGAGATATTIC	2000
ysGlyValLy	sHisLeuLys	AspMetProI	leLeuProG1	yGluIlePhe	
AAGTATAAAT	GGACAGTGAC	CGTAGAACAT	GGACCAACTA	AATCAGATOC	2050
LysTyrLysT	rpThrValTh	rValGluAsp	GlyProThrL	ysSerAspPr	
TOGGTGTGCTG	ACCOGATATT	ACTCAAGCTT	CATTAATCTG	GAGAGAGATC	2100
oArgCysLeu	ThrArgTyrT	yrSerSerPh	eIleAsnLeu	GluArgAspL	
TAGCTTCAGG	ACTCATTGGC	OCTCTCTCA	TCTGCTACAA	AGAATCTGIA	2150
euAlaSerG1	yLeuIleGly	ProLeuLeuI	leCysTyrLy	sGluSerVal	
GATCAAAGAG	GAAACCAGAT	GATGTACAGAC	AAGAGAAATG	TCATCCTGTT	2200
AspGlnArgG	lyAsnGlnMe	tMetSerAsp	LysArgAsnV	alIleLeuPh	
TTCIGTATT	GATGAGAATC	GAAGCTGGTA	CCTCACAGAG	AAATAIGCAGC	2250
eSerValPhe	AspGluAsnA	rgSerTrpTy	rLeuThrGlu	AsnMetGlnA	
GCTTCCTOOC	CAATGCAGAT	GTAGTGCAGC	CCCATGACCC	AGAGTTOCAA	2300
rgPheLeuPr	oAsnAlaAsp	ValValGlnP	roHisAspPr	oGluPheGln	
CTCTCTAAC	TCATGCACAG	CATCAATGGC	TATGTTTTG	ACAACCTGCA	2350
LeuSerAsnI	leMetHisSe	rIleAsnGly	TyrValPheA	spAsnLeuG1	
GCTGTCAGTT	TGTTTGCATG	AGGTGGCGTA	CTGGTACATT	CTAAGTGTIG	2400
nLeuSerVal	CysLeuHisG	luValAlaTy	rTrpTyrIle	LeuSerValG	
GAGCACAAAC	TGACTTCCTG	TCIGTCCTCT	TCCTCTGGATA	TACCTTCAAA	2450
lyAlaGlnTh	rAspPheLeu	SerValPheP	heSerGlyTy	rThrPheLys	
CACAAAATGG	TCTATGAAGA	CACACTTACC	CTCTTCCCCT	TCTCAGGAGA	2500
HisLysMetV	alTyrGluAs	pThrLeuThr	LeuPheProP	heSerGlyG1	
AACIGTCTTC	ATGTCAATGG	AAAACCCAGG	TCTGIGGGIT	CTGGGGTGCC	2550
uThrValPhe	MetSerMetG	luAsnProG1	yLeuTrpVal	LeuGlyCysH	
ACAACTCAGA	CITTCGGAAC	AGAGGCATGA	CAGCCTACT	GAAGGTTCT	2600
isAsnSerAs	pPheArgAsn	ArgGlyMetT	hrAlaLeuLe	uLysValSer	
AGITGTAAACA	GGAACATTGA	TGATTATTAT	GAGGACACAT	AOGAAGATAT	2650
SerCysAsnA	rgAsnIleAs	pAspTyrTyr	GluAspThrT	yrGluAspII	
TCCAACCTCCC	CTGCTAAATG	AAAACAATGT	AATTAAACCT	AGAAGCTCT	2700
eProThrPro	LeuLeuAsnG	luAsnAsnVa	lIleLysPro	ArgSerPheS	
CCCAGAATTIC	AAGGCACOCT	AGCACTAAGG	AAAAGCAATT	GAAAATGAAG	2750
erGlnAsnSe	rArgHisPro	SerThrLysG	luLysGlnLe	uLysMetLys	
AGAGAAGATT	TTGACATCTA	CGGCGACTAT	GAAAATCAGG	GCCTCCGGCAG	2800
ArgGluAspP	heAspIleTy	rGlyAspTyr	GluAsnGlnG	lyLeuArgSe	
CITTCAAAAG	AAAACACGAC	ACTATTCTAT	TGCTGCAGTG	GAGCGTCTCT	2850
rPheGlnLys	LysThrArgH	isTyrPheII	eAlaAlaVal	GluArgLeuT	

FIG. 6.D.

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10	20	30	40	50
1234567890	1234567890	1234567890	1234567890	1234567890
GGGATTATGG	GATGAGTAGA	TCTCCCCATA	TACTAAGAAA	CAGGGCTCAA
rpAspTyrGl	yMetSerArg	SerProHisI	leLeuArgAs	nArgAlaGln
AGTGGGGATG	TCCAGCAGTT	CAAGAAGGTG	GTTTCCAGG	AATTTACIGA
SerGlyAspV	alGlnGlnPh	eLysLysVal	ValPheGlnG	luPheThrAs
TGGATCCTT	ACTCAGOCCT	TATACCGTGG	AGAACTGAAT	GAACACTTGG
pGlySerPhe	ThrGlnProL	euTyrArgGl	yGluLeuAsn	GluHisLeuG
GACTCTGGG	GCATATATA	AGAGCAGAAG	TTGAAGACAA	TATCGTGGTA
lyLeuLeuGl	yProTyrIle	ArgAlaGluV	alGluAspAs	nIleValVal
ACTTCAAAAA	ACCAGGCCCTC	TOGTCCTAC	TCCTTCATT	CTAGCTTAT
ThrPheLysA	snGlnAlaSe	rArgProTyr	SerPheTyrS	erSerLeuII
TTCTATGAC	GAAGATGAGG	GACAAGGAGC	AGAACCTAGA	AGAAAGTTG
eSerTyrAsp	GluAspGluG	lyGlnGlyAl	aGluProArg	ArgLysPheV
TCAACCTAA	TGAAACCAAA	ATTACTTT	GGAAAGTGCA	GCATCATATG
alAsnProAs	nGluThrLys	IleTyrPheT	rpLysValGl	nHisHisMet
GCAACCACTA	AAGATGAGTT	TGACTGCAA	GCCTGGCTT	ATTTTCTGA
AlaProThrL	ysAspGluPh	eAspCysLys	AlaTrpAlaT	yrPheSerAs
TGTTGATTIG	GAGAAAGATG	TGCACTCAGG	CITGATTGGA	CCCCTCTGA
pValAspLeu	GluLysAspV	alHisSerGl	yLeuIleGly	ProLeuLeuI
TCTGCGCAG	TAACACACTG	AAACCTGCTC	ATGGGAGACA	AGTGACAGTG
1eCysArgSe	rAsnThrLeu	AsnProAlaH	isGlyArgGl	nValThrVal
CAGGAGTTG	CCCTGGTTT	CACTATATTIC	GATGAGACTA	AGAGCTGGTA
GlnGluPheA	1aLeuValPh	eThrIlePhe	AspGluThrL	ysSerTrpTy
CTTCACTGAA	AACCTGGAAA	GGAACTGTAG	AGCTCCCTGC	AATGTCCAGA
rPheThrGlu	AsnLeuGluA	rgAsnCysAr	gAlaProCys	AsnValGlnL
AGGAGGACCC	TACTCTAAA	GAAAACCTCC	GCTTCCATGC	AATCAAACGGC
ysGluAspPr	oThrLeuLys	GluAsnPheA	rgPheHisAl	ailleAsnGly
TATGTGAAGG	ATACACTCCC	TGGCTTAGTA	ATGGCTCAGG	ATCAAAAGGT
TyrValLysA	spThrLeuPr	oGlyLeuVal	MetAlaGlnA	spGlnLysVa
TOGATGGTAT	CTGCTCAGCA	TGGGCAGCAA	CGAAAACATT	CATTCCATTIC
1AngTrpTyr	LeuLeuSerM	etGlySerAs	nGluAsnIle	HisSerIleH
ACTTCAGTGG	ACATGTGTTTC	ACTGTACGGA	AAAAAGAGGA	ATATAAAATG
isPheSerGl	yHisValPhe	ThrValArgL	ysLysGluGl	uTyrLysMet
GCAGCTACA	ACCTCTATCC	AGGTGTTTT	GAGACTGTGG	AAATGCTACC
AlaValTyrA	snLeuTyrPr	oGlyValPhe	GluThrValG	luMetLeuPr
ATCCCAAGTT	GGAAATCTGGC	GGATAGAAATG	CCTTATCGGC	GAGCACCTGC
oSerGlnVal	GlyIleTrpA	rgIleGluCy	sLeuIleGly	GluHisLeuG
AAGCOGGGAT	GAGCACTCTG	TTTCTGGTGT	ACAGCAAGAA	GTGTCAAGACT
InAlaGlyMe	tSerThrLeu	PheLeuValT	yrSerLysLy	sCysGlnThr

FIG. 6.E.

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10	20	30	40	50
1234567890	1234567890	1234567890	1234567890	1234567890
CCACTGGGGA	TGGCTTGGG	ACACATTAGA	GATTTTCAGA	TTACAGCTTC
ProLeuGlyM	etAlaSerGl	yHisIleArg	AspPheGlnI	leThrAlaSe
AGGACAATAT	GGACAGTGCG	CCCCAAAGCT	GGCCAGACTT	CATTATTCOG
rGlyGlnTyr	GlyGlnTrpA	laProLysLe	uAlaArgLeu	HisTyrSerG
GATCAATCAA	TGCTTGGAGC	ACCAAGGATC	CCTTTTCTG	GATCAAGGIG
lySerIleAs	nAlaTrpSer	ThrLysAspP	roPheSerTr	pIleLysVal
GATCTCTTGG	CACCGATGAT	TATTCAGGC	ATCATGACCC	AGGGGGCCCG
AspLeuLeuA	laProMetI1	eIleHisGly	IleMetThrG	InGlyAlaAr
CCAGAACGTC	TOCAGGCTCT	ACGTGTCCTCA	GTTTATCATC	ATGTACAGTC
gGlnLysPhe	SerSerLeuT	yrValSerGl	nPheIleIle	MetTyrSerL
TGGATGGCAA	CAAGTGGCAC	AGTTACCGAG	GGAAITCCAC	GGGGAOCTTA
euAspGlyAs	nLysTrpHis	SerTyrArgG	lyAsnSerTh	rGlyThrLeu
ATGGCTCTCT	TTGGCAACGT	GGATTCACT	GGGATCAAAC	ACAATATTTC
MetValPheP	heGlyAsnVa	1AspSerSer	GlyIleLysH	isAsnIlePh
TAACOCTCOG	ATTATTGCTC	AGTACATOCG	TTTGCACCCA	ACCCATTACA
eAsnProPro	IleIleAlaG	InTyrIleAr	gLeuHisPro	ThrHisTyrS
GCATCOGCAG	CACTCTTCGC	ATGGAGCTCT	TGGGCTGTGA	CTTCAACAGT
erIleArgSe	rThrIleArg	MetGluLeuL	euGlyCysAs	pPheAsnSer
TGCAGCATGC	CGCTGGGGAT	GGAGAGTAAA	GCAATATCAG	ATGCTCAGAT
CysSerMetP	roLeuGlyMe	tGluSerLys	AlaIleSerA	spAlaGlnI1
CACTGCTCG	TCCTACCTAA	GCAGTATGCT	TGOCACCTTGG	TCTCCCTTCCC
eThrAlaSer	SerTyrLeuS	erSerMetLe	uAlaThrTrp	SerProSerG
AAGCCGGCT	GCACCTGCAG	GGCAGGACTA	ATGCTTGGAG	ACCTCAGGCA
InAlaArgLe	uHisLeuGln	GlyArgThrA	snAlaTrpAr	gProGlnAla
AATAACCAA	AAGAGTGGCT	GCAAGTGGAC	TTCCGGAAGA	CCATGAAAGT
AsnAsnProL	ysGluTrpLe	uGlnValAsp	PheArgLysT	hrMetLysVa
CACAGGAATA	ACCAACCAGG	GGGTGAAATC	TCTCTCATC	AGCATGTATG
lThrGlyIle	ThrThrGlnG	lyValLysSe	rLeuLeuIle	SerMetTyrV
TGAAGGAGTT	CCTCACCTCC	AGTAGTCAAG	ATGCCATAA	CTGGACTCTG
alLysGluPh	eLeuIleSer	SerSerGlnA	spGlyHisAs	nTrpThrLeu
TTCTTCAGA	ATGGCAAAGT	CAAGGCTCTC	CAGGGAAACC	GGGACTCCCTC
PheLeuGlnA	snGlyLysVa	1LysValPhe	GlnGlyAsnA	rgAspSerSe
CACGCTGTC	CGGAACCGTC	TCGAACCCCC	GCTGGTGGCT	CGCTAOGTGC
rThrProVal	ArgAsnArgL	euGluProPr	oLeuValAla	ArgTyrValA
GCCTGCACCC	GCAGAGCTGG	GCGCACCACA	TGCCCCCTGAG	GCTGGAGGGC
rgLeuHisPr	oGlnSerTrp	AlaHisHisI	leAlaLeuAr	gLeuGluVal
CTGGGCTGAG	ACACCCAGCA	GCGCGCTGA	CCCGCGCCCTC	TGCGGCGCTG
LeuGlyCysA	spThrGlnGl	nProAla...		

FIG. 6.F.

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10	20	30	40	50	
1234567890	1234567890	1234567890	1234567890	1234567890	
TCTCCCTGTC	CTCCCTGCGC	TGTCCCCGGG	GCTTCCCAC	AAGCTTATCG	4800
ATACCGTCTA	GCGAGTTCTT	CTGAGGGGAT	CGGCAATAAA	AAGACAGAAT	4850
AAAAAOGCAOG	GGTGTGGGT	GGTTGGTCTG	GATCCAGATC	TAGGAACOCC	4900
TAGTGATGGA	GTGGGOCACT	CCCTCTCTGC	GCGCTGGTC	GCTCACTGAG	4950
GGGGGGGGGG	CAAAGGGGGG	GCGTGGGGGG	ACCTTTGGTC	GGGGGGGCTC	5000
AGTGAGGAG	CGAGGAGGCA	GAGAGGGAGT	GGCCAAACCC	CCCCCCCCCC	5050
CCCCCTGCAGC	CCAGCTGCAT	TAATGAATCG	GCCAAOGCGC	GGGGAGAGGC	5100
GGTTGCGTA	TGGGGCGCTC	<u>TTCCGCTTCC</u>	<u>TCGCTCACIG</u>	<u>ACTCGCTGCG</u>	5150
<u>CTGGTCTT</u>	<u>GGCTGCGGC</u>	<u>GAGCGGTATC</u>	<u>AGCTCACICA</u>	<u>AAGGGGGTAA</u>	5200
<u>TACGGITATC</u>	<u>CACAGAATCA</u>	<u>GGGGATAACG</u>	<u>CAGGAAAGAA</u>	<u>CATGTGAGCA</u>	5250
<u>AAAGGCCACC</u>	<u>AAAAGGOCAG</u>	<u>GAACCGTAAA</u>	<u>AAGGCGCGT</u>	<u>TGCTGGCGTT</u>	5300
<u>TTTCCATAGG</u>	<u>CTCCGCCCCC</u>	<u>CTGACGAGCA</u>	<u>TCACAAAAAT</u>	<u>CGACGCTCAA</u>	5350
<u>GTCAGAGGTG</u>	<u>GCGAAACCCG</u>	<u>ACAGGACTAT</u>	<u>AAAGATACCA</u>	<u>GGCGTTTCCC</u>	5400
<u>CCTGGAAAGCT</u>	<u>CCCTCGTGGG</u>	<u>CTCTCCCTTT</u>	<u>COGACCCCTGC</u>	<u>CGCTTACCGG</u>	5450
<u>ATACCTGTC</u>	<u>GCCTTTCTCC</u>	<u>CTTGGGAAG</u>	<u>CGTGGCGCTT</u>	<u>TCTCAATGCT</u>	5500
<u>CAOGCTGTAG</u>	<u>GTATCTCAGT</u>	<u>TCGGTGTAGG</u>	<u>TOGTTGCTC</u>	<u>CAAGCTGGGC</u>	5550
<u>TGTGTGCACG</u>	<u>AACCCCCCGT</u>	<u>TCAGGCGAC</u>	<u>CGCTGOGCCT</u>	<u>TATCCGGTAA</u>	5600
<u>CTATCGCTT</u>	<u>GAGTCCAACC</u>	<u>CGGTAAGACA</u>	<u>CGACTTATCG</u>	<u>CCACTGGCAG</u>	5650
<u>CAGCCACTGG</u>	<u>TAACAGGATT</u>	<u>AGCAGAGGGA</u>	<u>GGTATGTAGG</u>	<u>GGGTGCTACA</u>	5700

FIG. 6.G.

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10	20	30	40	50	
1234567890	1234567890	1234567890	1234567890	1234567890	
<u>GAGTTCCTGA</u>	<u>AGTGGTGGGC</u>	<u>TAACTAACGC</u>	<u>TACACTAGAA</u>	<u>GGACAGTATT</u>	5750
<u>TGGTATCTGC</u>	<u>GCTCTGCTGA</u>	<u>AGCCAGTTAC</u>	<u>CTTOGGAAAA</u>	<u>AGAGTTGGTA</u>	5800
<u>GCTCTTGATC</u>	<u>CGGCAAACAA</u>	<u>ACCACOGCTG</u>	<u>GTAGCGGTGG</u>	<u>TTTTTTTGT</u>	5850
<u>TGCAAGCAGC</u>	<u>AGATTAACGCG</u>	<u>CAGAAAAAAA</u>	<u>GGATCTCAAG</u>	<u>AAGATCCCTT</u>	5900
<u>GATCTTTCT</u>	<u>ACGGGGTCTG</u>	<u>ACGCTCAGTG</u>	<u>GAACGAAAAC</u>	<u>TCACGTTAAG</u>	5950
<u>GGATTTTGGT</u>	<u>CATGAGAGTA</u>	<u>TCAAAAAGGA</u>	<u>TCTTCACCTA</u>	<u>GATCCTTTA</u>	6000
<u>AATTAAAAAT</u>	<u>GAAGTTTAA</u>	<u>ATCAATCTAA</u>	<u>AGTATATATG</u>	<u>AGTAAACTTG</u>	6050
<u>GTCIGACAGT</u>	<u>TACCAATGCT</u>	<u>TAATCAGTGA</u>	<u>GGCACCTATC</u>	<u>TCAGCGATCT</u>	6100
ylGelIreS ueL...siHo rPlaV...gr AueLreSgrA					
GTCIATTTCG TTCATCCATA GTTGCCTGAC TCCCCGTCGT GTAGATAACT					6150
psAelIulGn sAtelMprTue lnlGgrAlaV ylGgrAgrAr hTreSuel..					
AOGATAACGGG AGGGCTTACC ATCTGGCCCC AGTGTGCAA TGATACCGCG					6200
.reSlaVorP orPreSlaVt eMnlGylGpr TsihnlGueL reSlaValAu					
AGACOCCAOGC TCACCOGGCTC CAGATTIATC AGCAATAAAC CAGCCAGCOG					6250
eLy1GlaVre SlaVorPulG ueLnsAelIu eLueLueLyl GalAueLgrA					
GAAGGGGOOGA GOGCAGAAGT GGTCCTGCAA CTITATCCGC CTCCATCCAG					6300
ehPorPgrAa 1AsyCehPsi HpsAnlGueL syLeIigrAg rAprTylGrh					
TCTATTAATT GTTGCCTGGGA AGCTAGAGTA AGTAGITCGC CAGTTAATAG					6350
T.....nsA nsAylGorPu eL...ueLue LryThsAalA ueL...ryTn					
TTTGCCTAAC GTTGTGACA TIGCTACAGG CATOGTGGTG TCAOGCTCGT					6400
sAalAsyCgr AnlGnlGprT n1G...ueLs yCgrAorPrh TlaVreSrht					
CGTTTGGTAT GGCTTCATTC AGCTCOGGTT CCCAAOGATC AAGGCGAGTT					6450
rhTnlGryTo rPsyLteM... .reSgrAnsA ylGlaVelIu eLalAueL..					
ACATGATCCC CCATGTTGIG CAAAAAAAGCG GTTAGCTCCT TCGGTCTC					6500
.teMeliylG prTrhTrhTs yCehPueLor P...reSgrA grApsAulGr					
GATGTTGTC AGAAGTAAGT TGGCGCAGT GITATCACTC ATGGTTATGG					6550
eSgrAnsA... .ehPryTrhT orPgrAueLr hTelIlaV.. .orP...orP					
CAGCACTGCA TAATTCTCTT ACTGTCATGC CATCCGTAAG ATGCTTTCT					6600
ueLlaValAr yThsAulG... .n1G...alA teMgrAueLe 1IreSsyLnl					
GTGACTGGTG AGTACTCAAC CAAGTCATTG TGAGAATAGT GTATGOGGOG					6650
GreSnlGsiH rhTreSueLp rTrhTteMgr AueLelIrht ryTalAalAl					

FIG. 6.H.

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10	20	30	40	50	
<u>1234567890</u>	<u>1234567890</u>	<u>1234567890</u>	<u>1234567890</u>	<u>1234567890</u>	
AOCGAGITGC	TCTTGGCGGG	CGTCAATACG	GGATAATACC	GOGCACATA	6700
aVreSnsAre	SsyLylGorP	rhTueLlaVo	rPryTryTgr	AalAlaVryT	
GCAGAACTTT	AAAAGTGCTC	ATCATGGAA	AAOGITCTTC	GGGGCGAAAA	6750
syCehPsyLu	eLueLalA...n1GehP	1aVnsAsyLo	rPalAehPla	
CTCTCAAGGA	TCTTACCGCT	GTGAGATCC	AGITCGATGT	AAACCACTCG	6800
VgrAueLreS	grAlaValAr	hTreSelIpr	ThsAreSrht	1aVprTulGs	
TGCAACCAAC	TGATCTTCAG	CATCTTTAC	TTTCACCGAGC	GTTCCTGGGT	6850
iHlaVprTre	SelIsyLueL	teMsyL...s	yL...prTgr	AsyLn1GrhT	
GAGCAAAAC	AGGAAGGCAA	AATGOOGCAA	AAAAGGGAAT	AAGGGCGACA	6900
ueLueLehPu	eLehPalAeh	PsiHgrAueL	ehPorPehPu	eLorPreSla	
OGGAAATGTT	GAATACTCAT	ACTCTCCCT	TTTCAATATT	ATTGAAGCAT	6950
VreSelInsA	ehPlaV...	1			
TTATCAGGGT	TATTGCTCTCA	TGAGOGGATA	CATATTGAA	TGTATTAGA	7000
AAAATAAACAA	AATAGGGGTT	COGCGCACAT	TTCCCCGAAA	AGTGCCACCT	7050
GACGTCTAAG	AAACCATTAT	TATCATGACA	TTAACCTATA	AAAATAGGCG	7100
TATCACGAGG	CCCTTTCGTC	TCGCGCGTTT	CGGTGATGAC	GGTGAACACC	7150
TCTGACACAT	GCAGCTCCCG	GAGAOGGTCA	CAGCTTGTCT	GTAAGGGAT	7200
GGGGGAGCA	GACAAGCCCG	TCAGGGGOGOG	TCAGCGGGTG	TTGGCGGGTG	7250
TCGGGGCTGG	CTTAACATATG	CGGCATCAGA	GCAGATTGTA	CTGAGAGTGC	7300
ACCATATGCG	GTGTGAAATA	CGCACAGAT	GCGTAAGGAG	AAAATACCGC	7350
ATCAGGAAAT	<u>TGTAAACGTT</u>	<u>AATATTTGT</u>	<u>AAAAATTGCG</u>	<u>GTTAAATTGTT</u>	7400
<u>TGTAAATCA</u>	<u>GCTCATTTT</u>	<u>TAACCAATAG</u>	<u>GGCGAAATCG</u>	<u>GCAAAATCC</u>	7450
<u>TTATAAATCA</u>	<u>AAAGAATAGA</u>	<u>CGAGATAGG</u>	<u>GTGAGTGTT</u>	<u>GTTCAGITT</u>	7500
<u>GGAACAAGAG</u>	<u>TOCACTATTA</u>	<u>AAGAACGTGG</u>	<u>ACTCCAACGT</u>	<u>CAAAGGGCGA</u>	7550
<u>AAAACCGTCT</u>	<u>ATCAGGGCGA</u>	<u>TGGGCCACTA</u>	<u>CGTGAACCAT</u>	<u>CACCTAATC</u>	7600

FIG. 6.I.

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10	20	30	40	50	
1234567890	1234567890	1234567890	1234567890	1234567890	
<u>AAGT</u> TTTTTG	<u>GGGT</u> GAGGT	<u>GGGTAA</u> AGC	<u>ACTAAAT</u> CGG	<u>AAACCTAA</u> AG	7650
<u>GGAG</u> CCCCCG	<u>ATTA</u> AGAGCT	<u>TGAC</u> GGGGAA	<u>AGC</u> GGGCGA	<u>CGT</u> GGGAGA	7700
<u>AAGGAAGG</u> GA	<u>AGAAAG</u> CGAA	<u>AGGAG</u> GGGC	<u>GCTAGG</u> GC	<u>TGGCAAGI</u> GT	7750
<u>AGCGG</u> TCAOG	<u>CTG</u> OGOGTAA	<u>CCACCAC</u> ACC	<u>CGCG</u> GCGCTT	<u>AATG</u> GCGOOGC	7800
<u>TACAGGG</u> GC	<u>GTC</u> GGGOCAT	<u>TGOC</u> ATTCA	<u>GGCTACG</u> CAA	<u>CTG</u> TTGGAA	7850
<u>GGG</u> CGATCGG	<u>TGCG</u> GGCCTC	<u>TTCG</u> CATTAA	<u>CGC</u> AGCTGG	<u>CTG</u> CAGGGGG	7900
<u>GGGGGGGGGG</u>	<u>GGGT</u>				7914